

FIG. 1A

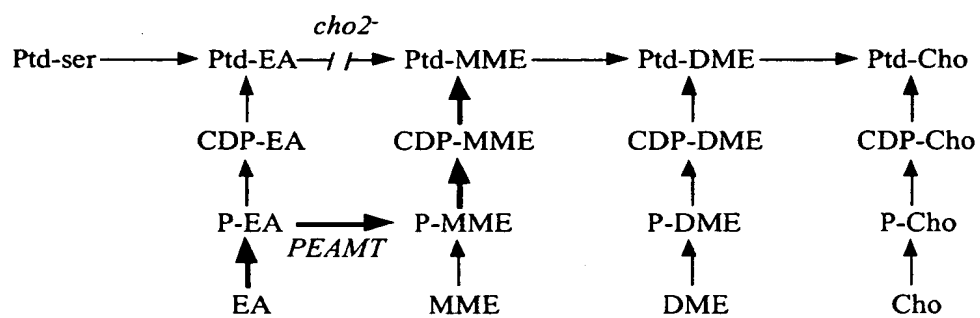


FIG. 1B

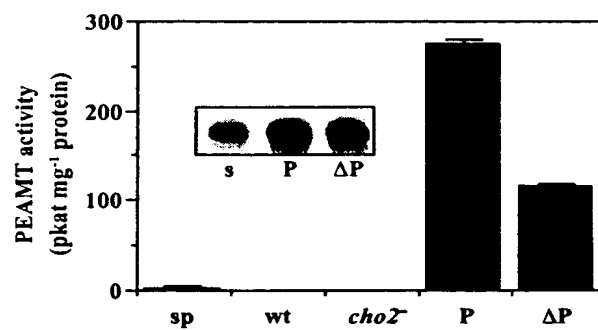


FIG. 2A

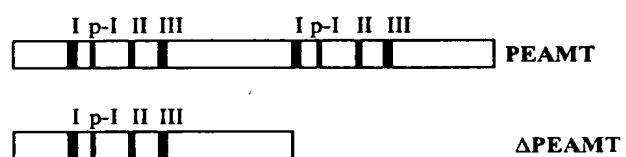


FIG. 2B

So 1 MAASAMCVLQERLVFKKYWIEHSVDLTVEAMMLDSQASDLDKVERPEVLSMLPPYEGKSVLELGAGIGRFTG
 Ac 1 MAASFMEREETRDIQKNYWIEHSADLTVEAMMLDSRASDLDKVERPEVLSMLPPYEGKSVLELGAGIGRFTG

post-I II

So 73 ELAKASQVIALDFIHSVIKKNESINGHYKNVKFMCADVTSFSLNIEPNSVDIIFSNWLLMYLSDKEEVERLV
 Ac 73 ELAKAGELIALDFIDNVIKKNESINGHYKNVKFMCADVTSPLKIIDGSLDLIFSNWLLMYLSDKEVELLA

III

So 145 ERMKKWLKPGGGYIFFRESCFHQSGLHKKRKSNGTHYREPRFYTKIFKECHMQDSDGNSIELSLIGCKCIGAYV
 Ac 145 ERMVGIKVGGGYIFFRESCFHQSGLSKRKSNPTHYREPRFYKVFQECQTRDAAGNSIELSLIGCKCIGAYV

I post-I II

So 217 KSKKNQNOISWLVQKVDSDDIGFORFLDSSQYKFNSTLRYERVFGPGYVSTGGLETTKEFVSKLDLKPGQK
 Ac 217 KMKKNQNOICWLVQKVSSENDRGFORFLDNVOYKSSGILRYERVFGQGTVSTGGLETTKEFVEKMLKPGQK

I post-I II

So 289 VLDVGGCGIGGGDFYMAENYDVHVVGIDLSINMISFALERSIGLKCAREFEVADCTKKDYFINSFDVIYSRDT
 Ac 289 VLDVGGCGIGGGDFYMAEKFDVHVVGIDLSVNMISFALERAIGLSCSVEFEVADCTTKHYPDNSFDVIYSRDT

III

So 361 ILHIQDKPALFRSFHKKWLKPGGKVLISDYCKSACTPSAEFAAYIRQGYDLHDVKAYGKMLKDAGFVEVIAE
 Ac 361 ILHIQDKPALFRITFFKKWLKPGGKVLISDYCRSPKTPSAEFSRYIKQGYDLHDVQAYGQMLKDAGFTDVIAE

So 433 NRTDQFIQVLLQKELDALEQEKDDFIDDFSIEDYNDIVDGKKAKLVPTTEGECQWGLFIAKKM
 Ac 433 DRTDQFMQVLLKRELDREVEKEKEKFIIDDFSIEDYNDIVDGKKAKLVPTTEGECQWGLFIAKKM

FIG. 3

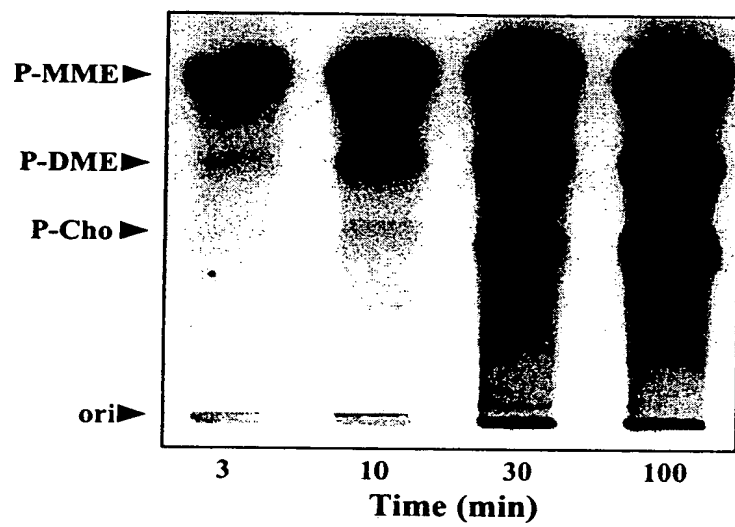


FIG. 4A

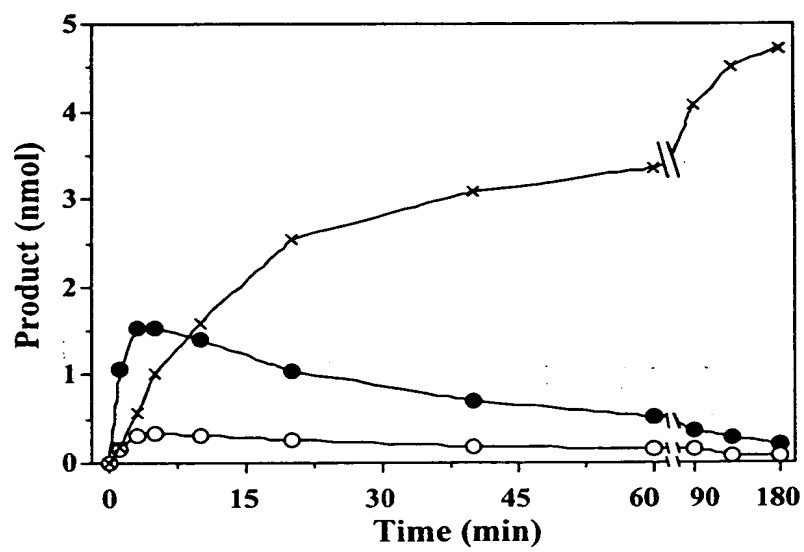


FIG. 4B

096363.034500

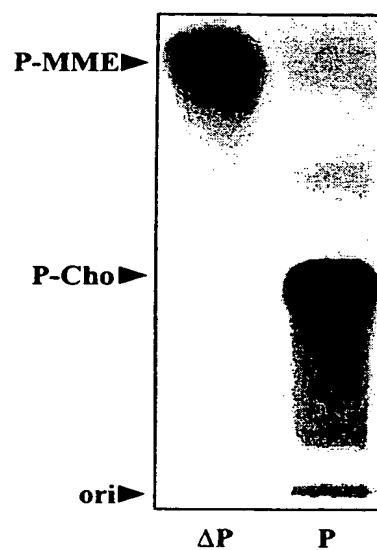


FIG. 5A

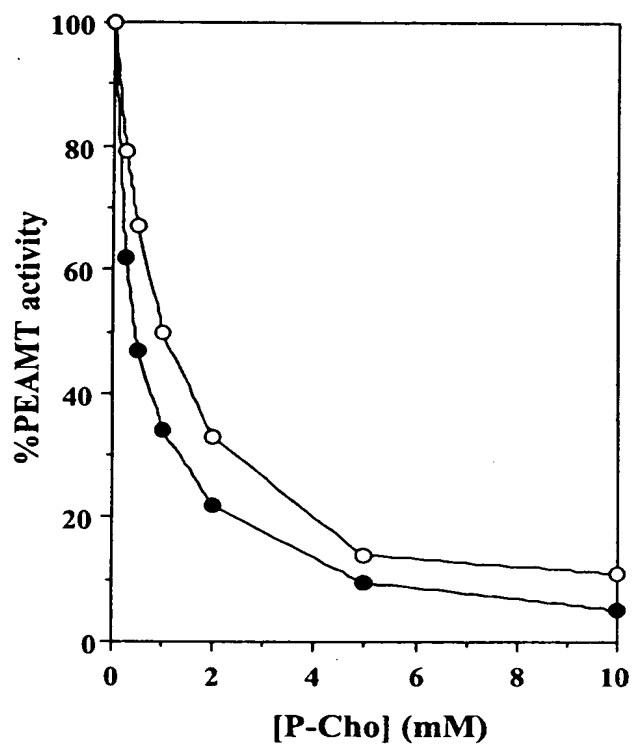


FIG. 5B

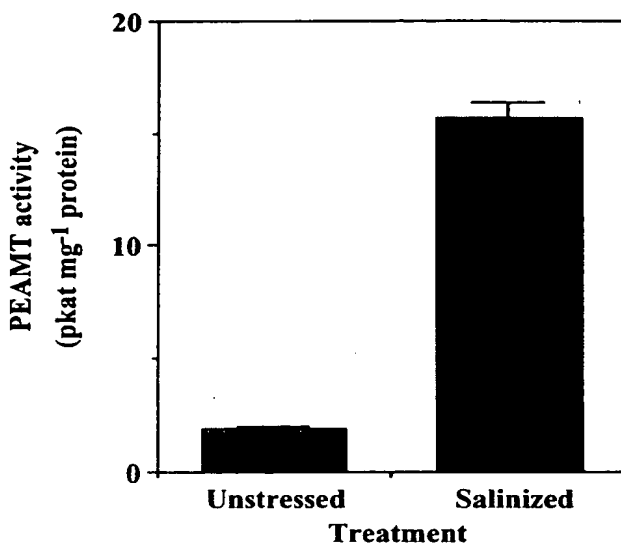


FIG. 6